



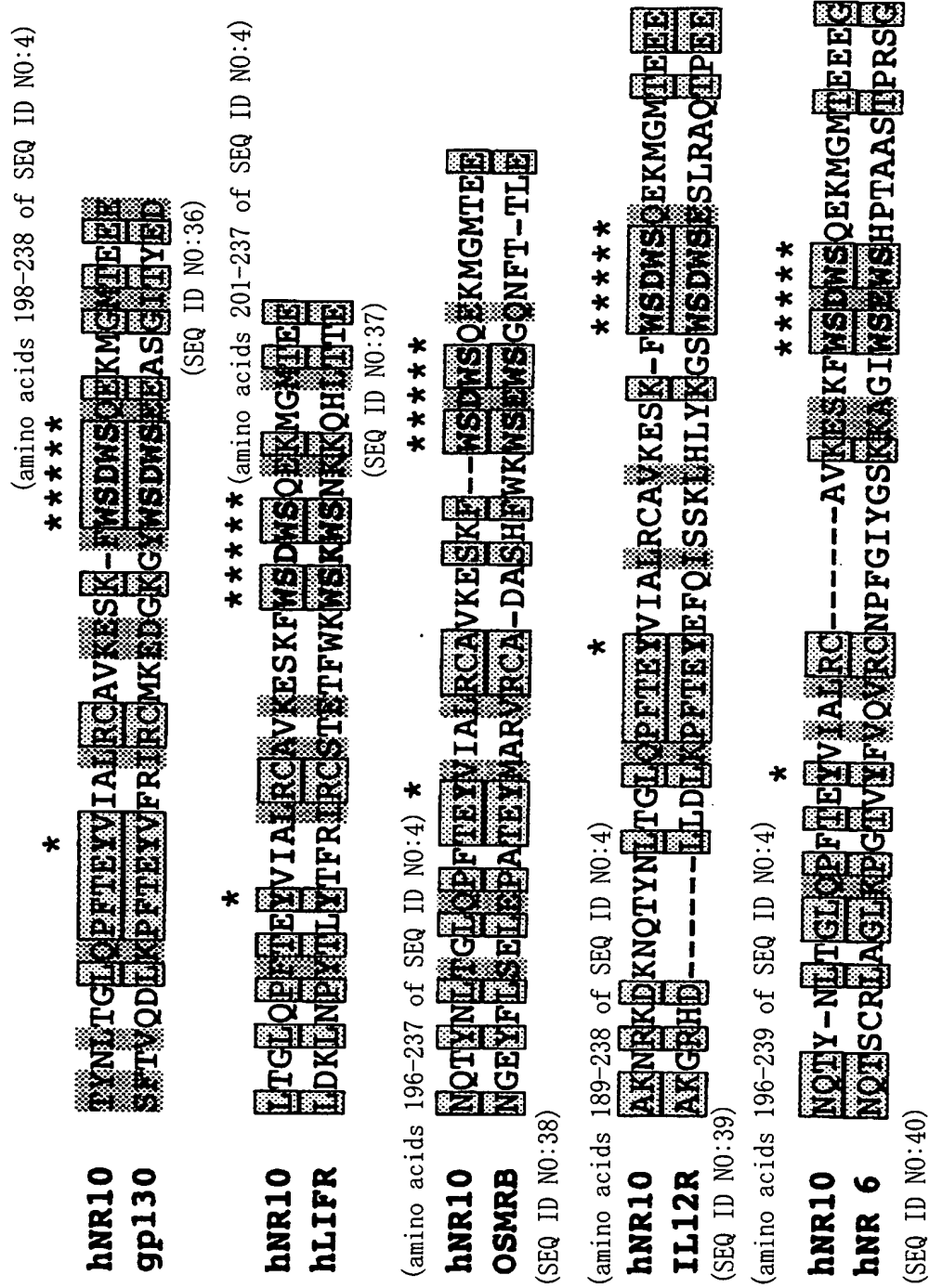
Figure 1

1 ttgggtggttcattggtgatgttctatctgtgttaagtaccaattgttcccaggcacatat  
61 ggaagtctgttaataaaaaatgatataattttaaaaatttgatttagagtgttactagttcta  
121 aaaatgtaaaagtacactaggtagtgaagaggaaaaatgggaggataaacgtgtgggtctcca  
181 ttccagtttacgattgtctctgtcttgtagatggaagtcaacttcgctaagaaccgtaag  
MetGluValAsnPheAlaLysAsnArgLys  
241 gataaaaaaccacacgtacaacctcacggggctgcaacctctttacagaatatgtcatagct  
AspLysAsnGlnThrTyrAsnLeuThrGlyLeuGlnProXxxThrGluTyrValIleAla  
301 ctgcgatgtgcgggtcaaggagtcacaaagttctggagtgactggagccaagaaaaaatggga  
LeuArgCysAlaValLysGluSerLysPheTrpSerAspTrpSerGlnGluLysMetGly  
361 atgactgaggaagaaagcaagctacttccctgcgattcccgtcctgtctgctctgggtgtan  
MetThrGluGluGluXxxLysLeuLeuProAlaIlePro (SEQ ID NO:35)  
421 ggctgctctgcgotaaaacttgggtgggtgtctgcaccacgg (SEQ ID NO:34)



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Figure 2





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Figure 3

(SEQ ID NO:1)

1 CGCTTATAAATGAATGTGTGCTTAGGAACACCAGACAGCACTCCAGCACTCTGCTTGGGG  
61 GGCATTTCGAAACAGCAAAATCACTCATAAAAGGCAAAAAATTGCAAAAAAATAGTAATA  
121 ACCAGCATGGTACTAAATAGACCATGAAAAGACATGTGTGTGCAGTATGAAAATTGAGAC  
181 AGGAAGGCAGAGTGTCTGCTTCCACCTCAGCTGGGAATGTGCATCAGGCAACTCAAG  
241 TTTTTCACCACGGCATGTGTCTGTGAATGTCCGCAAAACATTTTAACAATAATGCAATCC  
301 ATTTCCCAGCATAAGTGGGTAAGTGCCACTTTGACTTGGGCTGGGCTTAAAGCACAAGA  
361 AAAGCTCGCAGACAATCAGAGTGGAAACACTCCACATCTTAGTGTGGATAAATTAAAGT  
421 CCAGATTGTTCTTCTGTCTGACTTGTGCTGTGGGAGGTGGAGTTGCCTTTGATGCAAA  
481 TCCTTTGAGCCAGCAGAACATCTGTGGAACATCCCCTGATACATGAAGCTCTCTCCCCAG

(SEQ ID NO:2) MetLysLeuSerProGln

541 CCTTCATGTGTTAACCTGGGGATGATGTGGACCTGGGCACTGTGGATGCTCCCTCACTC  
ProSerCysValAsnLeuGlyMetMetTrpThrTrpAlaLeuTrpMetLeuProSerLeu  
601 TGCAAATTCAGCCTGGCAGCTCTGCCAGCTAAGCCTGAGAACATTTCTGTGTCTACTAC  
CysLysPheSerLeuAlaAlaLeuProAlaLysProGluAsnIleSerCysValTyrTyr  
661 TATAGGAAAAATTTAACCTGCACTTGGAGTCCAGGAAAGGAAACCAGTTATACCCAGTAC  
TyrArgLysAsnLeuThrCysThrTrpSerProGlyLysGluThrSerTyrThrGlnTyr  
721 ACAGTTAAGAGAACTTACGCTTTCGGAGAAAAACATGATAATTGTACAACCAATAGTTCT  
ThrValLysArgThrTyrAlaPheGlyGluLysHisAspAsnCysThrThrAsnSerSer  
781 ACAAGTGAATTCGTGCTTCGTGCTCTTTTTTCTTCCAAGAATAACGATCCAGATAAT  
ThrSerGluAsnArgAlaSerCysSerPhePheLeuProArgIleThrIleProAspAsn  
841 TATACCATTGAGGTGGAAGCTGAAAATGGAGATGGTGTAAATTAAATCTCATATGACATAC  
TyrThrIleGluValGluAlaGluAsnGlyAspGlyValIleLysSerHisMetThrTyr  
901 TGGAGATTAGAGAACATAGCGAAACTGAACCACCTAAGATTTTCCGTGTGAAACCAGTT  
TrpArgLeuGluAsnIleAlaLysThrGluProProLysIlePheArgValLysProVal  
961 TTGGGCATCAAACGAATGATTCAAATGAATGGATAAAGCCTGAGTTGGCGCTGTTTCA  
LeuGlyIleLysArgMetIleGlnIleGluTrpIleLysProGluLeuAlaProValSer  
1021 TCTGATTTAAATACACACTTCGATTCAGGACAGTCAACAGTACCAGCTGGATGGAAGTC  
SerAspLeuLysTyrThrLeuArgPheArgThrValAsnSerThrSerTrpMetGluVal  
1081 AACTTCGCTAAGAACCCTAAGGATAAAACCAAACGTACAACCTCACGGGGCTGCAGCCT  
AsnPheAlaLysAsnArgLysAspLysAsnGlnThrTyrAsnLeuThrGlyLeuGlnPro  
1141 TTTACAGAATATGTCATAGCTCTGCGATGTGCGGTCAAGGAGTCAAAGTCTCGAGTGC  
PheThrGluTyrValIleAlaLeuArgCysAlaValLysGluSerLysPheTrpSerAsp



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Figure 4

(SEQ ID NO:1)

1201 TGGAGCCAAGAAAAAATGGGAATGACTGAGGAAGAAGCTCCATGTGGCCTGGAAGTGTGG

(SEQ ID NO:2)

TrpSerGlnGluLysMetGlyMetThrGluGluGluAlaProCysGlyLeuGluLeuTrp

1261 AGAGTCCTGAAACCAGCTGAGCGGATGGAAGAAGCCAGTGCAGTTGTTATGGAAGAAG

ArgValLeuLysProAlaGluAlaAspGlyArgArgProValArgLeuLeuTrpLysLys

1321 GCAAGAGGAGCCCCAGTCTAGAGAAAACACTTGGCTACAACATATGGTACTATCCAGAA

AlaArgGlyAlaProValLeuGluLysThrLeuGlyTyrAsnIleTrpTyrTyrProGlu

1381 AGCAACACTAACCTCACAGAAACAATGAACACTACTAACCAGCAGCTTGAAGTGCATCTG

SerAsnThrAsnLeuThrGluThrMetAsnThrThrAsnGlnGlnLeuGluLeuHisLeu

1441 GGAGGCGAGAGCTTTTGGGTGTCTATGATTCTTATAATTCTCTTGGGAAGTCTCCAGTG

GlyGlyGluSerPheTrpValSerMetIleSerTyrAsnSerLeuGlyLysSerProVal

1501 GCCACCCTGAGGATTCCAGCTATTCAAGAAAAATCATTTCAAGTGCATTGAGGTCATGCAG

AlaThrLeuArgIleProAlaIleGlnGluLysSerPheGlnCysIleGluValMetGln

1561 GCCTGCGTTGCTGAGGACCAGCTAGTGGTGAAGTGGCAAAGCTCTGCTCTAGACGTGAAC

AlaCysValAlaGluAspGlnLeuValValLysTrpGlnSerSerAlaLeuAspValAsn

1621 ACTTGGATGATTGAATGGTTTCCGGATGTGGACTCAGAGCCCACCACCCTTTCTGGGAA

ThrTrpMetIleGluTrpPheProAspValAspSerGluProThrThrLeuSerTrpGlu

1681 TCTGTGTCTCAGGCCACGAACTGGACGATCCAGCAAGATAAATTAACCTTTCTGGTGC

SerValSerGlnAlaThrAsnTrpThrIleGlnGlnAspLysLeuLysProPheTrpCys

1741 TATAACATCTCTGTGTATCCAATGTTGCATGACAAAGTTGGCGAGCCATATCCATCCAG

TyrAsnIleSerValTyrProMetLeuHisAspLysValGlyGluProTyrSerIleGln

1801 GCTTATGCCAAAGAAGGCGTTCCATCAGAAGGTCTGAGACCAAGGTGGAGAACATTGGC

AlaTyrAlaLysGluGlyValProSerGluGlyProGluThrLysValGluAsnIleGly

1861 GTGAAGACGGTCACGATCACATGGAAGAGATTCCCAAGAGTGAGAGAAAGGGTATCATC

ValLysThrValThrIleThrTrpLysGluIleProLysSerGluArgLysGlyIleIle

1921 TGCAACTACACCATCTTTACCAAGCTGAAGGTGGAAGGATTCTCCAAGACAGTCAAT

CysAsnTyrThrIlePheTyrGlnAlaGluGlyGlyLysGlyPheSerLysThrValAsn

1981 TCCAGCATCTTGAGTACGGCCTGGAGTCCCTGAAACGAAAGACCTCTTACATTGTTTAC

SerSerIleLeuGlnTyrGlyLeuGluSerLeuLysArgLysThrSerTyrIleValGln

2041 GTCATGGCCAACACCAAGTGTGGGGGAACCAACGGGACCAGCATAAATTTCAAGACATTG

ValMetAlaAsnThrSerAlaGlyGlyThrAsnGlyThrSerIleAsnPheLysThrLeu

2101 TCATTCAAGTGTCTTTGAGATTATCCTCATAACTTCTCTGATTGGTGGAGGCCTTCTTATT

SerPheSerValPheGluIleIleLeuIleThrSerLeuIleGlyGlyGlyLeuLeuIle



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Figure 5

2161 CTCATTATCCTGACAGTGGCATATGGTCTCAAAAAACCCAACAAATTGACTCATCTGTGT  
LeuIleIleLeuThrValAlaTyrGlyLeuLysLysProAsnLysLeuThrHisLeuCys  
2221 TGGCCACCGTCCCAACCCTGCTGAAAGTAGTATAGCCACATGGCATGGAGATGATTTC  
TrpProThrValProAsnProAlaGluSerSerIleAlaThrTrpHisGlyAspAspPhe  
2281 AAGGATAAGCTAAACCTGAAGGAGTCTGATGACTCTGTGAACACAGAAGACAGGATCTTA  
LysAspLysLeuAsnLeuLysGluSerAspAspSerValAsnThrGluAspArgIleLeu  
2341 AAACCATGTTCCACCCCAGTGACAAGTTGGTGATTGACAAGTTGGTGGTGAACCTTTGGG  
LysProCysSerThrProSerAspLysLeuValIleAspLysLeuValValAsnPheGly  
2401 AATGTTCTGCAAGAAATTTTCACAGATGAAGCCAGAACGGGTCAGGAAAAACAATTTAGG  
AsnValLeuGlnGluIlePheThrAspGluAlaArgThrGlyGlnGluLysGlnPheArg  
2461 AGGGGAAAAGAATGGGACTAGAATTCTGTCTTCCTGCCCAACTTCAATATAAGTGTGGAC  
ArgGlyLysGluTrpAsp\*\*\* (SEQ ID NO:2)  
2521 TAAAATGCGAGAAAGGTGTCCTGTGGTCTATGCAAATTAGAAAGGACATGCAGAGTTTTTC  
2581 CAACTAGGAAGACTGAATCTGTGGCCCCAAGAGAACCATCTCCGAAGACTGGGTATGTGG  
2641 TCTTTTCCACACATGGACCACCTACGGATGCAATCTGTAATGCATGTGCATGAGAAGTCT  
2701 GTTATTAAGTAGAGTGTGAAAACATGGTTATGGTAATAGGAACAGCTTTTAAAATGCTTT  
2761 TGTATTTGGGCCTTTCACACAAAAAAGCCATAATACCATTTTCATGTAATGCTATACTTC  
2821 TATACTATTTTCATGTAATACTATACTTCTATACTATTTTCATGTAATACTATACTTCTA  
2881 TACTATTTTCATGTAATACTATACTTCTATATTAAAGTTTTACCCACTCCAAAAAAGAA  
2941 AAAAAAAAAAAAAAAAAAAAAAAAAAAAAA (SEQ ID NO:1)



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Figure 6

(SEQ ID NO:3)

1 CGCTTATAAATGAATGTGTGCTTAGGAACACCAGACAGCACTCCAGCACTCTGCTTGGGG  
61 GGCATTTCGAAACAGCAAATCACTCATAAAAGGCAAAAATTGCAAAAAAATAGTAATA  
121 ACCAGCATGGTACTAAATAGACCATGAAAAGACATGTGTGTGCAGTATGAAAATTGAGAC  
181 AGGAAGGCAGAGTGTCTAGCTTGTCCACCTCAGCTGGGAATGTGCATCAGGCAACTCAAG  
241 TTTTTCACCAACGGCATGTGTCTGTGAATGTCCGCAAAACATTTTAAACAATAATGCAATCC  
301 ATTTCCAGCATAAGTGGGTAAGTGCCACTTTGACTTGGGCTGGGCTTAAAAGCACAAGA  
361 AAAGCTCGCAGACAATCAGAGTGGAAACACTCCACATCTTAGTGTGGATAAATTAAAGT  
421 CCAGATTGTTCTTCTGTCTGACTTGTGTGTGGGAGGTGGAGTTGCCTTTGATGCAAA  
481 TCCTTTGAGCCAGCAGAACATCTGTGGAACATCCCCTGATACATGAAGCTCTCTCCCCAG

(SEQ ID NO:4) MetLysLeuSerProGln

541 CCTTCATGTGTTAACCTGGGGATGATGTGGACCTGGGCACTGTGGATGCTCCCTCACTC  
ProSerCysValAsnLeuGlyMetMetTrpThrTrpAlaLeuTrpMetLeuProSerLeu  
601 TGCAAATTCAGCCTGGCAGCTCTGCCAGCTAAGCCTGAGAACATTTCTGTGTCTACTAC  
CysLysPheSerLeuAlaAlaLeuProAlaLysProGluAsnIleSerCysValTyrTyr  
661 TATAGGAAAAATTTAACCTGCACTTGGAGTCCAGGAAAGGAAACCAGTTATACCCAGTAC  
TyrArgLysAsnLeuThrCysThrTrpSerProGlyLysGluThrSerTyrThrGlnTyr  
721 ACAGTTAAGAGAACTTACGCTTTCGGAGAAAAACATGATAATTGTACAACCAATAGTTCT  
ThrValLysArgThrTyrAlaPheGlyGluLysHisAspAsnCysThrThrAsnSerSer  
781 ACAAGTGAAAAATCGTGCTTCGTGCTCTTTTTCCTTCCAAGAATAACGATCCAGATAAT  
ThrSerGluAsnArgAlaSerCysSerPhePheLeuProArgIleThrIleProAspAsn  
841 TATACCATTGAGGTGGAAGCTGAAAATGGAGATGGTGTAAATTAAATCTCATATGACATAC  
TyrThrIleGluValGluAlaGluAsnGlyAspGlyValIleLysSerHisMetThrTyr  
901 TGGAGATTAGAGAACATAGCGAAACTGAACCACCTAAGATTTTCCGTGTGAAACCAGTT  
TrpArgLeuGluAsnIleAlaLysThrGluProProLysIlePheArgValLysProVal  
961 TTGGGCATCAAACGAATGATTCAAATTGAATGGATAAAGCCTGAGTTGGCGCCTGTTTCA  
LeuGlyIleLysArgMetIleGlnIleGluTrpIleLysProGluLeuAlaProValSer  
1021 TCTGATTTAAAATACACACTTCGATTTCAGGACAGTCAACAGTACCAGCTGGATGGAAGTC  
SerAspLeuLysTyrThrLeuArgPheArgThrValAsnSerThrSerTrpMetGluVal  
1081 AACTTCGCTAAGAACCGTAAGGATAAAAACCAACGTACAACCTCACGGGGCTGCAGCCT  
AsnPheAlaLysAsnArgLysAspLysAsnGlnThrTyrAsnLeuThrGlyLeuGlnPro  
1141 TTTACAGAATATGTCATAGCTCTGCGATGTGCGGTCAAGGAGTCAAAGTTCTGGAGTGAC  
PheThrGluTyrValIleAlaLeuArgCysAlaValLysGluSerLysPheTrpSerAsp



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Figure 7

1201 TGGAGCCAAGAAAAAATGGGAATGACTGAGGAAGAAGGCAAGCTACTCCCTGCGATTCCC  
TrpSerGlnGluLysMetGlyMetThrGluGluGluGlyLysLeuLeuProAlaIlePro  
1261 GTCCTGTCTACTCTGGTGTAGGGCTGCTTTGGGCTAGACTTGGTGGGGTTTGTCAACCACC  
ValLeuSerThrLeuVal\*\*\* (SEQ ID NO:4)

1321 TGGTTGGGAATCATGGAATCTCATGACCCAGGGGCCCCCTGTACCATCGAGAGTGAGCC  
1381 TGCACAACCTTTGTGCCCCAAAGGCAAAGGATCACATTTTAATACTCATGAGGTTCTTATA  
1441 CTATACATGAAAGGGTATCATATCATTGTGTTTGTGTTTGTGTTTGTGTTTGTGAGATGGAGTC  
1501 TTA CTCTGTCAACCAGGATGGAGTGCAGTGATGTGATCTCGGCTCACTGCCACCACCACC  
1561 TCCCGAGTTCAAGCAATTCTTGTGCCTCAGCCTCCCAAGTAGCTGGGATTACAGGGGCCC  
1621 ACGACCATGCCCCGGTTGATTTTTGTATTTTTAGTAGAGAAGGGATATCACCATGTTGGCT  
1681 AGGCTAGTCTTGAACTCCTGACCTCAGGTAATCTGCCCACCTTGACCTCCCAAAGTGTG  
1741 GGATTACAGGCGTGAGCCACTGTGCCCGCCAGTATCATATCATCTGAAGGTATCCTGTG  
1801 ATAAATTAAAGATACATATTGTGAATCCTGGAGCTACTACTCAAAAAATAAATAAAGGTG  
1861 TAACTAATACAATTTAAAAAATCACATTTTAAATGACAGTGAGGAAAGGAAAGAGGCATG  
1921 GATTGCAGGTTGATGGAGTGCTTACTAAGTGTCAGTATGGTCATTAAGAGCAACGCTTCC  
1981 AGTCAGTGGCCTTGGCTTAAATCCCAAGCCAGGTGTCTTTGGGCAAGATACCTAAACTCT  
2041 CAGTTCATTCTCAGCAGTTTCCTCGCATTTATTCCCCTTTTCTATATTGAAATAGAATAT  
2101 GTAAGTTGAGTTTATAGTAGTACCTATTTTTTAGTATTATTTTAAAGATTAAATGAAATA  
2161 ATGTGTTTAGCCCATAGTAGATATTCCTAACTGCTAGACTTCCTATTCTTATTATTAT  
2221 CCTCCTACTATTATTTTAAATCCTCCTTAAAGCACTATAAAATATGTAGAGTCACTCCCA  
2281 TTTTGGAAATGAGGAACTGAGTTTCAGAGATGCTAATAAACAGCTCAGGGTCACTCAGC  
2341 ATGTGTTACTTTTCTCAAGAGCCTTGCCAGAGTCTGACCCCTCAGTGGACGATCAATAAA  
2401 TGTGTGATGAATGGAAAAAAAAAAAAAAAAAAAAAAAAAAAA (SEQ ID NO:3)



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Figure 13

(SEQ ID NO:16)

1 CCCCTGATACATGAAGCTCTCTCCCCAGCCTTCATGTGTAACTGGGGATGATGTGGAC

(SEQ ID NO:17) MetLysLeuSerProGlnProSerCysValAsnLeuGlyMetMetTrpThr

61 CTGGGCACTGTGGATGCTCCCTCACTCTGCAAATTCAGCCTGGCAGCTCTGCCAGCTAA  
TrpAlaLeuTrpMetLeuProSerLeuCysLysPheSerLeuAlaAlaLeuProAlaLys

121 GCCTGAGAACATTTCTGTGTCTACTACTATAGGAAAAATTTAACCTGCACCTGGAGTCC  
ProGluAsnIleSerCysValTyrTyrTyrArgLysAsnLeuThrCysThrTrpSerPro

181 AGGAAAGGAAACCAGTTATACCCAGTACACAGTTAAGAGAACTTACGCTTTTGGAGAAAA  
GlyLysGluThrSerTyrThrGlnTyrThrValLysArgThrTyrAlaPheGlyGluLys

241 ACATGATAATTGTACAACCAATAGTTCTACAAGTGAAAAATCGTGCTTCGTGCTCTTTTTT  
HisAspAsnCysThrThrAsnSerSerThrSerGluAsnArgAlaSerCysSerPhePhe

301 CCTTCCAAGAATAACGATCCAGATAATTATACCATTTGAGGTGGAAGCTGAAAAATGGAGA  
LeuProArgIleThrIleProAspAsnTyrThrIleGluValGluAlaGluAsnGlyAsp

361 TGGTGTAATTAAATCTCATATGACATACTGGAGATTAGAGAACATAGCGAAACTGAACC  
GlyValIleLysSerHisMetThrTyrTrpArgLeuGluAsnIleAlaLysThrGluPro

421 ACCTAAGATTTTCCGTGTGAAACCAGTTTGGGCATCAAACGAATGATTCAAATTTGAATG  
ProLysIlePheArgValLysProValLeuGlyIleLysArgMetIleGlnIleGluTrp

481 GATAAAGCCTGAGTTGGCGCCTGTTTCATCTGATTTAAAATACACACTTCGATTCAGGAC  
IleLysProGluLeuAlaProValSerSerAspLeuLysTyrThrLeuArgPheArgThr

541 AGTCAACAGTACCAGCTGGATGGAAGTCAACTTCGCTAAGAACCGTAAGGATAAAAAACCA  
ValAsnSerThrSerTrpMetGluValAsnPheAlaLysAsnArgLysAspLysAsnGln

601 AACGTACAACCTCACGGGGCTGCAGCCTTTTACAGAATATGTCATAGCTCTGCGATGTGC  
ThrTyrAsnLeuThrGlyLeuGlnProPheThrGluTyrValIleAlaLeuArgCysAla

661 GGTCAAGGAGTCAAAGTTCTGGAGTGACTGGAGCCAAGAAAAATGGGAATGACTGAGGA  
ValLysGluSerLysPheTrpSerAspTrpSerGlnGluLysMetGlyMetThrGluGlu

721 AGAAGCTCCATGTGGCCTGGAAGTGTGGAGAGTCTGAAACCAGCTGAGGCGGATGGAAG  
GluAlaProCysGlyLeuGluLeuTrpArgValLeuLysProAlaGluAlaAspGlyArg

781 AAGGCCAGTGCAGTTGTTATGGAAGAAGCAAGAGGAGCCCCAGTCCTAGAGAAAACACT  
ArgProValArgLeuLeuTrpLysLysAlaArgGlyAlaProValLeuGluLysThrLeu

841 TGGCTACAACATATGGTACTATCCAGAAAGCAACACTAACCTCACAGAAACAATGAACAC  
GlyTyrAsnIleTrpTyrTyrProGluSerAsnThrAsnLeuThrGluThrMetAsnThr

901 TACTAACCAGCAGCTTGAAGTGCATCTGGGAGGCGAGAGCTTTTGGGTGTCTATGATTTT  
ThrAsnGlnGlnLeuGluLeuHisLeuGlyGlyGluSerPheTrpValSerMetIleSer

961 TTATAATTCTCTTGGGAAGTCTCCAGTGGCCACCCTGAGGATTCCAGCTATTCAAGAAAA  
TyrAsnSerLeuGlyLysSerProValAlaThrLeuArgIleProAlaIleGlnGluLys

1021 ATCATTTTCAGTGCATTGAGGTCATGCAGGCCTGCGTTGCTGAGGACCAGCTAGTGGTGAA





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Figure 14

SerPheGlnCysIleGluValMetGlnAlaCysValAlaGluAspGlnLeuValValLys  
1081 GTGGCAAAGCTCTGCTCTAGACGTGAACACTTGGATGATTGAATGGTTTCCGGATGTGGA  
TrpGlnSerSerAlaLeuAspValAsnThrTrpMetIleGluTrpPheProAspValAsp  
1141 CTCAGAGCCCACCACCCTTTCTGGGAATCTGTGTCTCAGGCCACGAACCTGGACGATCCA  
SerGluProThrThrLeuSerTrpGluSerValSerGlnAlaThrAsnTrpThrIleGln  
1201 GCAAGATAAATTAAACCTTTCTGGTGCTATAACATCTCTGTGTATCCAATGTTGCATGA  
GlnAspLysLeuLysProPheTrpCysTyrAsnIleSerValTyrProMetLeuHisAsp  
1261 CAAAGTTGGCGAGCCATATTCCATCCAGGCTTATGCCAAAGAAGGCGTTCCATCAGAAGG  
LysValGlyGluProTyrSerIleGlnAlaTyrAlaLysGluGlyValProSerGluGly  
1321 TCCTGAGACCAAGGTGGAGAACATTGGCGTGAAGACGGTCACGATCACATGGAAGAGAT  
ProGluThrLysValGluAsnIleGlyValLysThrValThrIleThrTrpLysGluIle  
1381 TCCCAAGAGTGAGAGAAAGGGTATCATCTGCAACTACACCATCTTTTACCAAGCTGAAGG  
ProLysSerGluArgLysGlyIleIleCysAsnTyrThrIlePheTyrGlnAlaGluGly  
1441 TGGAAAAGGATTCTCCAAGACAGTCAATTCCAGCATCTTGCAAGTACGGCCTGGAGTCCCT  
GlyLysGlyPheSerLysThrValAsnSerSerIleLeuGlnTyrGlyLeuGluSerLeu  
1501 GAAACGAAAGACCTCTTACATTGTTCAAGTCATGGCCAGCACCAGTCTGGGGGAACCAA  
LysArgLysThrSerTyrIleValGlnValMetAlaSerThrSerAlaGlyGlyThrAsn  
1561 CGGGACCAGCATAAATTTCAAGACATTGTCATTCAAGTGTCTTTGAGATTATCCTCATAAC  
GlyThrSerIleAsnPheLysThrLeuSerPheSerValPheGluIleIleLeuIleThr  
1621 TTCTCTGATTGGTGGAGGCCTTCTTATTCTCATTATCCTGACAGTGGCATATGGTCTCAA  
SerLeuIleGlyGlyGlyLeuLeuIleLeuIleIleLeuThrValAlaTyrGlyLeuLys  
1681 AAAACCCAAACAAATTGACTCATCTGTGTTGGCCCACCGTTCCCAACCCTGCTGAAAGTAG  
LysProAsnLysLeuThrHisLeuCysTrpProThrValProAsnProAlaGluSerSer  
1741 TATAGCCACATGGCATGGAGATGATTCAAGGATAAGCTAAACCTGAAGGAGTCTGATGA  
IleAlaThrTrpHisGlyAspAspPheLysAspLysLeuAsnLeuLysGluSerAspAsp  
1801 CTCTGTGAACACAGAAGACAGGATCTTAAACCATGTTCCACCCCCAGTGACAAGTTGGT  
SerValAsnThrGluAspArgIleLeuLysProCysSerThrProSerAspLysLeuVal  
1861 GATTGACAAGTTGGTGGTGAACCTTTGGGAATGTTCTGCAAGAAATTTTCACAGATGAAGC  
IleAspLysLeuValValAsnPheGlyAsnValLeuGlnGluIlePheThrAspGluAla  
1921 CAGAACGGGTCAAGAAAACAATTTAGGAGGGGAAAAGAATGGGACTAGAATTCTGTCTTC  
ArgThrGlyGlnGluAsnAsnLeuGlyGlyGluLysAsnGlyThrArgIleLeuSerSer  
1981 CTGCCCAACTTCAATATAAGTGTGGACTAAAATGCGAGAAAGGTGTCTGTGGTCTATGC  
CysProThrSerIle\*\*\* (SEQ ID NO:17)  
2041 AAATTAGAAAGGACATGCAGAGTTTTTCCAAC TAGGAAGACTGAATCTGTGGCCCCAAGAG  
2101 AACCATCTCCGAAGACTGG (SEQ ID NO:16)